
Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: [year=2008; month=5; day=7; hr=13; min=58; sec=37; ms=649;]

Reviewer Comments:

<210> 53

<211> 48

<212> DNA

<213> Artificial

<220>

<223> Primer (seq. 1 and 2) or Aptamer (all others)

<220>

<221> misc_feature

<222> (1)..(1)

<223> n is 0 to 1000 of any nucleotides or spacer molecules

<220>

<221> misc_feature

<222> (8)..(8)

<223> n is 0 to 1000 of any nucleotides or spacer molecules

<220>

<221> misc_feature

<222> (13)..(13)

<223> n is 0 to 1000 of any nucleotides or spacer molecules

Invalid response for 'n' locations as shown in the above attachment.
'n' represent only single nucleotide. Please check for similar errors in subsequent sequences.

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<220>
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<221> misc_feature

<222> (44)..(44)

<223> n is u or bind

'n' has a valid response as u but bind is not acceptable, as 'n' respresent single nucleotide. check for similar errors in subsequent sequences.

Validated By CRFValidator v 1.0.3

Application No: 10527345 Version No: 1.0

Input Set:

Output Set:

Started: 2008-04-24 16:29:25.550

Finished: 2008-04-24 16:29:27.101

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 551 ms

Total Warnings: 72

Total Errors: 0

No. of SeqIDs Defined: 72

Actual SeqID Count: 72

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Input Set:

Output Set:

Started: 2008-04-24 16:29:25.550 **Finished:** 2008-04-24 16:29:27.101

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 551 ms

Total Warnings: 72
Total Errors: 0

No. of SeqIDs Defined: 72

Actual SeqID Count: 72

Error code Error Description

This error has occured more than 20 times, will not be displayed

SEQUENCE LISTING

<110>	RiNA Netzwerk RNA-Technologien GmbH Rimmele, Martina Orgel, Dagmar
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<130>	7003/35
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<141>	2008-04-24
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99

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98

98

9

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<223> Primer (seq. 1 and 2) or Aptamer (all others)

<223> Primer (seq. 1 and 2) or Aptamer (all others)

<223> Primer (seq. 1 and 2) or Aptamer (all others)

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ugccggguuc accaugaacg aacguagcuu cccuauga

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                                                                      97
ucgcagucca cgucgguccu cgggcauuug ugccccc
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ugccaguccg gcgguugcua uaguaggagu ggguuauagu
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                                                                     102
uccgccuuau ugccauuuga auaacuugug cgcgaguaca ca
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aacuucaacg gacgauggug ugcggcuggg accccaugcg ugc
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<223> Primer (seq. 1 and 2) or Aptamer (all others)

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cccucauugg cccncgaccu ggcacagcaa agcancauag g
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auauuuugag cccauguuga gaaacggugc cccacuuaaa gc
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<223> n is a, c, g, t or u
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                                                                      98
ugacunuuac auuccagcag cucgaccuag cggcgccc
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<223> Primer (seq. 1 and 2) or Aptamer (all others)
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gccggcuaac ggaugucccu acgcaugacc ugcauucacc g
                                                                    101
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<211> 48
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<220>
<223> Primer (seq. 1 and 2) or Aptamer (all others)
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<222> (14)..(14)
<223> n is g or c
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<223> Primer (seq. 1 and 2) or Aptamer (all others)

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<220>
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